RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/529, 749
Source:	TFWO
Date Processed by STIC:	09/08/2006
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IFWO

RAW SEQUENCE LISTING DATE: 09/08/2006
PATENT APPLICATION: US/10/529,749 TIME: 08:55:25

Input Set : A:\10529749.txt

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3 <110> APPLICANT: Japan Science and Technology Agency
        KURODA, Shunichi
 4
         TANIZAWA, Katsuyuki
 5
         KONDO, Akihiko
 6
 7
         UEDA, Masakazu
 8
         SENO, Masaharu
10 <120> TITLE OF INVENTION: Hollow Protein Nanoparticles, and Drug Using the Same
12 <130> FILE REFERENCE: 1035-579 / P023P08US
14 <140> CURRENT APPLICATION NUMBER: US 10/529,749
15 <141> CURRENT FILING DATE: 2005-03-30
17 <150> PRIOR APPLICATION NUMBER: PCT/JP2003/015003
18 <151> PRIOR FILING DATE: 2003-11-25
20 <150> PRIOR APPLICATION NUMBER: JP 2002-339925
21 <151> PRIOR FILING DATE: 2002-11-22
23 <160> NUMBER OF SEQ ID NOS: 6
25 <170> SOFTWARE: MS Word
27 <210> SEQ ID NO: 1
28 <211> LENGTH: 1218
29 <212> TYPE: DNA
30 <213> ORGANISM: Hepatitis B virus
32 <220> FEATURE:
33 <221> NAME/KEY: CDS
34 <222> LOCATION: (1)..(1218)
36 <400> SEQUENCE: 1
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38 Met Arg Ser Leu Leu Ile Leu Val Leu Cys Phe Leu Pro Leu Ala Ala
                     5
                                        10
41 ttg ggt aag gtt cga caa ggc atg ggg acg aat ctt tct gtt ccc aat
42 Leu Gly Lys Val Arg Gln Gly Met Gly Thr Asn Leu Ser Val Pro Asn
43
                20
                                    25
45 cct ctg gga ttc ttt ccc gat cac cag ttg gac cct gcg ttc gga gcc
                                                                      144
46 Pro Leu Gly Phe Phe Pro Asp His Gln Leu Asp Pro Ala Phe Gly Ala
49 aac tca aac aat cca gat tgg gac ttc aac ccc aac aag gat caa tgg
                                                                      192
50 Asn Ser Asn Asn Pro Asp Trp Asp Phe Asn Pro Asn Lys Asp Gln Trp
51
        50
                            55
53 cca gag gca aat cag gta gga gcg gga gca ttc ggg cca ggg ttc acc
                                                                      240
54 Pro Glu-Ala Asn Gln Val Gly Ala Gly Ala Phe Gly Pro Gly Phe Thr
                        70
                                             75
57 cca cca cac ggc ggt ctt ttg ggg tgg agc cct cag gct cag ggc ata
                                                                      288
58 Pro Pro His Gly Gly Leu Leu Gly Trp Ser Pro Gln Ala Gln Gly Ile
                                        90
61 ttg aca aca gtg cca gca gca cct cct cct gcc tcc acc aat cgg cag
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PATENT APPLICATION: US/10/529,749 TIME: 08:55:25

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	Leu	Thr	Tnr	Val	Pro	Ala	Ala	Pro		Pro	Ala	ser	Thr		Arg	GIn	
63	+ a a	~~~	200	100		20+			105					110			204
				cag Gln													384
67	261	GIY	115	GIII	PIO	TIIL	PIO	120	ser	PIO	PIO	ьец	125	Asp	ser	піѕ	
	aat	cac		a to	a=a	taa	22t		202	202	tta	a aa		~a+	ata	at a	422
		_	_	atg Met	_									_	_		432
71	PIO	130	AIG	Met	GIII	пр	135	ser	TILL	1111	Pne	140	GIII	нта	ьеu	ьеu	
	gat		202	gtg	200	aaa		tat	+++	aat	aat		~~~	+ ~ ~	20t	taa	480
				Val													400
	145	110	****9	vai	nr 9	150	шси	- 7 -	1110	110	155	Gry	Gry	Der	Ser	160	
		aca	ata	aac	cct		cca	act	act	acc		CCC	ata	tot	aaa		528
				Asn													520
79	017		• • • •	11011	165	• • • •	110			170	DCI	110	110	DCI	175	нър	
-	cat	gca	cca	aac		gag	aac	aca	aca		gga	ttc	cta	gga		cta	576
				Asn	-	-										_	• • •
83				180					185		0-1			190		_	
	ctc	ata	tta	cag	aca	aaa	ttt	ttc		tta	aca	aga	atc		aca	ata	624
				Gln					_	_		_					
87			195			-		200	•			,	205				
89	cca	cag	agt	cta	gac	tcg	tgg	tgg	act	tct	ctc	aat	ttt	cta	ggg	gga	672
		_	_	Leu	_	_											
91		210			-		215	-				220			-	-	
93	gca	CCC	acg	tgt	cct	ggc	caa	aat	tcg	cag	tcc	cca	acc	tcc	aat	cac	720
94	Ala	Pro	Thr	Cys	Pro	Gly	Gln	Asn	Ser	Gln	Ser	Pro	Thr	Ser	Asn	His	
95	225					230					235					240	
97	tca	cca	acc	tct	tgt	cct	cca	att	tgt	cct	ggc	tat	cgc	tgg	atg	tgt	768
98	Ser	Pro	Thr	Ser	Cys	Pro	Pro	Ile	Cys	${\tt Pro}$	Gly	Tyr	Arg	Trp	Met	Cys	
99					245					250					255		
																c atc	816
		ı Arç	g Arg			: Ile	Phe	. Let	ı Phe	e Ile	e Lei	ı Let	ı Leı	ı Cys	s Lei	ı Ile	
103				260					265					270	-		
																cct	864
		e Lei			. Let	ı Let	ı Asp	_		ı Gly	/ Met	: Let			L Cys	s Pro	
107			275					280					285				
																tgc	912
				o GIŽ	Thr	Ser			Sei	Thi	C GI		_	з Гуз	Thi	r Cys	
111		290					295		- •			300				•	
																aca	960
			Pro	O Ala	GIT	_		Sei	. Met	: Phe			Cys	s Cys	s Cys	s Thr	
	305					310					315					320	
			_	_			_		_							tgg	1008
	_	Pro	sei	c Asp	_		Cys	Thr	Cys) ITE	Pro	Sei		Trp	
119					325					330					335		
																tgg	1056
		, Lue	: ATG			: ьeu	LIT	GIU			ı sei	. val	. Arg			r Trp	
123		~ -	. +	340					345					350			7704
																C CCC	1104
126	, тел	ı ser	. net	т тел	val	. Pro	, FUE	· val	. GII	ıırı	, FUE	e val	. GI	тer	ı sei	r Pro	

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127			355					360					365				
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	Thr																
131		370					375					380		1			
	ctg		aac	atc	tta	agt		ttt	tta	cct	cta		cca	att	ttc	ttt	1200
	Leu				_	_											
	385	-1-				390			200		395	200				400	
	tgt	ctt	taa	ata	tat						333					100	1218
	Cys			-													1210
139	CID	200		• • •	405												
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	4 <212> TYPE: PRT																
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149		9	DCI	ЦСЦ	5	110	DCu	vui	LCu	10	THE	шец	110	шец	15	AIG	
	Leu	Glv	Lvc	Val		Gln	Glv	Met	Glv		Δen	T.011	Sor	Val		Δen	
152	D Cu	Oly	Lys	20	n. 9	GIII	Gry	NCC	25	1111	ASII	пси	Ser	30	110	ASII	
	Pro	T.eu	Glv		Dhe	Pro	Aen	Hic		T.e.11	Δen	Dro	Δla		Glv	Δla	
155	110	шси	35	1110	1110	110	пор	40	GIII	neu	иор	110	45	rnc	Cly	AIG	
	Asn	Ser	-	Δsn	Pro	Asn	Trn		Dhe	Δen	Dro	Δen		Δen	Gln	מאנה	
158		50			110	1101	55	1105	1110	11011	110	60	_	тор	0111		
	Pro		Δla	Asn	Gln	Val		Δla	Glv	Δla	Phe		Pro	Glv	Phe	Thr	
161		O.L.		11011	0111	70	017		017	1114	75	O _T	110	O _T	1110	80	
	Pro	Pro	His	Glv	Glv		Len	Glv	Trp	Ser		Gln	Ala	Gln	Glv		
164					85			U -1		90		0111		0111	95		
	Leu	Thr	Thr	Val		Ala	Δla	Pro	Pro		Δla	Ser	Thr	Asn		Gln	
167				100					105					110	5		
	Ser	Glv	Ara		Pro	Thr	Pro	Tle		Pro	Pro	Leu	Ara		Ser	His	
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173		130					135					140					
175	Asp	Pro	Arq	Val	Arq	Gly		Tyr	Phe	Pro	Ala	Gly	Gly	Ser	Ser	Ser	
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178	Gly	Thr	Val	Asn	Pro	Val	Pro	Thr	Thr	Ala	Ser	Pro	Ile	Ser	Gly	Asp	
179	•				165					170					175	-	
181	Pro	Ala	Pro	Asn	Met	Glu	Asn	Thr	Thr	Ser	Gly	Phe	Leu	Gly	Pro	Leu	
182				180					185		•			190			
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187	Pro	Gln	Ser	Leu	Asp	Ser	Trp	Trp	Thr	Ser	Leu	Asn	Phe	Leu	Gly	Gly	
188		210			_		215	_				220			-	•	•
190	Ala	Pro	Thr	Cys	Pro	Gly	Gln	Asn	Ser	Gln	Ser	Pro	Thr	Ser	Asn	His	
	225			-		230					235					240	
	Ser	Pro	Thr	Ser	Cys	Pro	Pro	Ile	Cys	Pro		Tyr	Arg	Trp	Met		
194					245				_	250	_	_		•	255	-	
196	Leu	Arg	Arg	Phe	Ile	Ile	Phe	Leu	Phe	Ile	Leu	Leu	Leu	Cys	Leu	Ile	
197		-	_	260					265					270			

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Output Set: N:\CRF4\09082006\J529749.raw

199 Phe Leu Leu Val Leu Leu Asp Tyr Gln Gly Met Leu Pro Val Cys Pro 200 275 280 202 Leu Leu Pro Gly Thr Ser Thr Ser Thr Gly Pro Cys Lys Thr Cys 295 205 Thr Ile Pro Ala Gln Gly Thr Ser Met Phe Pro Ser Cys Cys Cys Thr 206 305 310 315 208 Lys Pro Ser Asp Gly Asn Cys Thr Cys Ile Pro Ile Pro Ser Ser Trp 325 330 211 Ala Phe Ala Arg Phe Leu Trp Glu Trp Ala Ser Val Arg Phe Ser Trp 345 214 Leu Ser Leu Leu Val Pro Phe Val Gln Trp Phe Val Gly Leu Ser Pro 355 360 217 Thr Val Trp Leu Ser Val Ile Trp Met Met Trp Tyr Trp Gly Pro Ser 375 220 Leu Tyr Asn Ile Leu Ser Pro Phe Leu Pro Leu Pro Ile Phe Phe 221 385 390 395 223 Cys Leu Trp Val Tyr Ile 227 <210> SEQ ID NO: 3 228 <211> LENGTH: 27 229 <212> TYPE: DNA 230 <213> ORGANISM: Artificial Sequence 232 <220> FEATURE: 233 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer 235 <400> SEQUENCE: 3 236 gccggtaccg cgagcttacc agttctc 27 239 <210> SEQ ID NO: 4 240 <211> LENGTH: 27 241 <212> TYPE: DNA 242 <213> ORGANISM: Artificial Sequence 244 <220> FEATURE: 245 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer 247 <400> SEQUENCE: 4 248 gccctcgagg aaactaagtt tcttggt 27 251 <210> SEQ ID NO: 5 252 <211> LENGTH: 30 253 <212> TYPE: DNA 254 <213> ORGANISM: Artificial Sequence 256 <220> FEATURE: 257 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer 259 <400> SEQUENCE: 5 260 gccctcgaga ccatggacat tgacccgtat 30 263 <210> SEQ ID NO: 6 264 <211> LENGTH: 39 265 <212> TYPE: DNA 266 <213> ORGANISM: Artificial Sequence 268 <220> FEATURE: 269 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer 271 <400> SEQUENCE: 6

RAW SEQUENCE LISTING

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Input Set : A:\10529749.txt

Output Set: N:\CRF4\09082006\J529749.raw

272 gccgagctcc taaacgcgtc cacattgaga ttcccgaga

39

VERIFICATION SUMMARY

DATE: 09/08/2006

PATENT APPLICATION: US/10/529,749

TIME: 08:55:26

Input Set : A:\10529749.txt